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TOPOLOGY: both

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Ser Arg Asp Arg	Gln His Leu Pro Leu	Ile Lys Met Lys Pro	Val
200	205	210	
Pro Asp Leu Val	Pro Gly Asn Phe Lys	Ser Gln Leu Gln Lys	Val
215	220	225	
Pro Pro Glu Trp	Lys Ala Leu Thr Asp	Met Pro Gln Met Arg	Met
230	235	240	
Glu Leu Glu Arg	Pro Gly Gly Asn Glu	Ile Thr Arg Gly Gly	Ser
245	250	255	
Thr Ser Tyr Gly	Thr Gly Ser Glu Thr	Glu Ser Pro Arg Asn	Pro
260	265	270	
Ser Ser Ala Gly	Ser Trp Asn Ser Gly	Ser Ser Gly Pro Gly	Ser
275	280	285	
Thr Gly Asn Arg	Asn Pro Gly Ser Ser	Gly Thr Gly Gly Thr	Ala
290	295	300	
Thr Trp Lys Pro	Gly Ser Ser Gly Pro	Gly Ser Thr Gly Ser	Trp
305	310	315	
Asn Ser Gly Ser	Ser Gly Thr Gly Ser	Thr Gly Asn Gln Asn	Pro
320	325	330	
Gly Ser Pro Arg	Pro Gly Ser Thr Gly	Thr Trp Asn Pro Gly	Ser
335	340	345	
Ser Glu Arg Gly	Ser Ala Gly His Trp	Thr Ser Glu Ser Ser	Val
350	355	360	
Ser Gly Ser Thr	Gly Gln Trp His Ser	Glu Ser Gly Ser Phe	Arg
365	370	375	
Pro Asp Ser Pro	Gly Ser Gly Asn Ala	Arg Pro Asn Asn Pro	Asp
380	385	390	
Trp Gly Thr Phe	Glu Glu Val Ser Gly	Asn Val Ser Pro Gly	Thr
395	400	405	
Arg Arg Glu Tyr	His Thr Glu Lys Leu	Val Thr Ser Lys Gly	Asp
410	415	420	
Lys Glu Leu Arg	Thr Gly Lys Glu Lys	Val Thr Ser Gly Ser	Thr
425	430	435	
Thr Thr Thr Arg	Arg Ser Cys Ser Lys	Thr Val Thr Lys Thr	Val
440	445	450	
Ile Gly Pro Asp	Gly His Lys Glu Val	Thr Lys Glu Val Val	Thr
455	460	465	
Ser Glu Asp Gly	Ser Asp Cys Pro Glu	Ala Met Asp Leu Gly	Thr
470	475	480	
Leu Ser Gly Ile	Gly Thr Leu Asp Gly	Phe Arg His Arg His	Pro
485	490	495	

Asp	Glu	Ala	Ala	Phe	Phe	Asp	Thr	Ala	Ser	Thr	Gly	Lys	Thr	Phe	
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Pro	Gly	Phe	Phe	Ser	Pro	Met	Leu	Gly	Glu	Phe	Val	Ser	Glu	Thr	
				515					520					525	
Glu	Ser	Arg	Gly	Ser	Glu	Ser	Gly	Ile	Phe	Thr	Asn	Thr	Lys	Glu	
				530					535					540	
Ser	Ser	Ser	His	His	Pro	Gly	Ile	Ala	Glu	Phe	Pro	Ser	Arg	Gly	
				545					550					555	
Lys	Ser	Ser	Ser	Tyr	Ser	Lys	Gln	Phe	Thr	Ser	Ser	Thr	Ser	Tyr	
				560					565					570	
Asn	Arg	Gly	Asp	Ser	Thr	Phe	Glu	Ser	Lys	Ser	Tyr	Lys	Met	Ala	
				575					580					585	
Asp	Glu	Ala	Gly	Ser	Glu	Ala	Asp	His	Glu	Gly	Thr	His	Ser	Thr	
				590					595					600	
Lys	Arg	Gly	His	Ala	Lys	Ser	Arg	Pro	Val	Arg	Gly	Ile	His	Thr	
				605					610					615	
Ser	Pho	Leu	Gly	Lys											
				620											

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 <213> Artificial Sequence
 <220>
 <221>
 <222>
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 STRANDEDNESS: single
 TOPOLOGY: linear
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 Gly Pro Arg Pro